Sequence Listing

SEQ ID NO. 1

Q04984 and AAH23518
Chaperonin 10

```
 1 maggafrkfl plfdrvlver saaetvtkgg imlpeksqgk vlqatvvavg sgskgkggei
61 qpvsvkvgdk vllpeyggtk vvlddkdyfl frdgdilgky vd
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SEQ ID NO. 2

NM_002157 and U07550
Human chaperonin 10 mRNA, complete cds

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  1 gctacactag agcagagtac gagtctgagg cggagggagt aatggcagga caagcgttta
 61 gaaagtttct tccactcttt gaccgagtat tggttgaaag gagtgctgct gaaactgtaa
121 ccaaaggagg cattatgctt ccagaaaaat ctcaaggaaa agtattgcaa gcaacagtag
181 tcgctgttgg atcgggttct aaaggaaagg gtggagagat tcaaccagtt agcgtgaaag
241 ttggagataa agttcttctc ccagaatatg gaggcaccaa agtagttcta gatgacaagg
301 attatttcct atttagagat ggtgacattc ttggaaagta cgtagactga aataagtcac
361 tattgaaatg gcatcaacat gatgctgccc attccactga agttctgaaa tctttcgtca
421 tgtaaataat ttccatattt ctcttttata ataaactaat gataactaat gacatccagt
481 gtctccaaaa ttgtttcctt gtactgatat aaacacttcc aaataaaaat atgtaaat
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SEQ ID NO. 3

P05109
Calgranulin A

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 1 mltelekaln siidvyhkys likgnfhavy rddlkkllet ecpqyirkkg advwfkeldi
61 ntdgavnfqe flilvikmgv aahkkshees hke
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SEQ ID NO. 4

A12027
Macrophage migration inhibition factor (MRP-14)cDNA from Human placenta (formula v)

```
  1 cttgggttgc ttccaccttt tggctcttgt aaataatgct gctatgaaca tgaatgtaca
 61 aacatctgtt tgaatccctg cattcaattc ttttgcatat atacccagga gcagaatgat
121 ggatcatatg gtaattctgt gtttatttat ttgaggaaca aacttgccgt tttccataac
181 agctgcacta ttttacattc ccactaacag tgcattaggc ttccaattct ctatgccctc
241 accaacactt gttttctggg ttttaaaaga agtagtagtc atccttgtag gtgtcaggtg
301 gtatctcatt gtcgttttgc ttcatgtttt cctaaagatt agtaattttc atatgcttat
361 tgaccatttg tatatcttct tcggagaagt gtctatttga gtctttcccc aattttgatt
421 ggtttgtttg ttttttgttg ttgagttgta gggattcttt tatattctgg atattaatcc
481 cttatcagat atttgtttta caaatatttt ctttgtaaca acagaaacac accacagtct
541 tcaaggttgg aagccagtta atctgagtag cattttgtta gtggtgggga gaggatttgt
601 tcctcctgaa atcctgggga attggccacc tcctcttctc ctcttaggca tgaagcgcgt
661 ctggcttctc caaagaactc ttcccctcca ctacctcaga gttagcttcc tctcttcagc
721 cagtgatcct ggggtcccag acacaataat taaccaagag agggtgaaag gctccctgct
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781 gtgtttatgc aatggctcag gcccttgtga agtgccgagg gaccccaagc agcctccatc
841 tcccagggca tggtccatcc ccagctttca cagaacagga aagctgtgga ggagtgtggg
901 cagcagggta ggaatggata tagcccttgg caacaacaca tttccccaca aagcacccac
961 ccaaaagaac aacaacgata gttttagttt ttagtaatga gaacaatagt tctcatgact
1021 aaaagccatc agccaggaca ctgttctcaa ccctttttgcg gtctttggac cctttgaaac
1081 tctgacagaa gccatggagg aatgttctca ctgagtgcat gcactcaaaa tgatgcattc
1141 aacttcaatt cagtttcagg gatgtatggc ctgaccacca atgcagggga ttagcaatcg
1201 caatagtgga gagggcatgg gagtgggaat ctggctggat caagcaagtg gatgccagca
1261 gcccagaaaa agagcccccc tacctgcttt ttccttcctg ggcactattg cccagcaaat
1321 gccttcctct ttccgcttct cctacctccc cacccaaaat tttcattctg cacagtgatt
1381 gccacattca ctggttgaga aacagagact gtagcaactc tggcagggag aagctgtctc
1441 tgatggcctg aagctgtggg cagctggcca agcctaaccg ctataaaaag gagctgcctc
1501 tcagccctgc atgtctcttg tcagctgtct ttcagaagac ctggtaagtg ggactgtctg
1561 ggttggcccc gcactttggg cttctcttgg ggagggtcag ggaagtggag cagccttcct
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1681 tgaggcaggg ataaggaagg agagtatcct ccagcacctt ccagtgggta agggcacatt
1741 gtctcctagg ctggactttt cttgagcaga gggtggggtg gtaaggaaag tctacgggcc
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1861 ctatcatccc acccttccca ccagaggcca tagccatctg ctggtttggt tatttgagag
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1981 atgcaaattc cctgccatgg gattccccag aaggttctgt ttttcaggtg gggcaagttc
2041 cgtgggcatc atgttgaccg agctggagaa agccttgaac tctatcatcg acgtctacca
2101 caagtactcc ctgataaagg ggaatttcca tgccgtctac agggatgacc tgaagaaatt
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2281 ctatgcctcg gctctctctg agatctttaa actctggctt cttcctcctc aatcttgaca
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2401 cttccaggag ttcctcattc tggtgataaa gatgggcgtg gcagcccaca aaaaaagcca
2461 tgaagaaagc cacaaagagt agctgagtta ctgggcccag aggctgggcc cctggacatg
2521 tacctgcaga ataataaagt catcaatacc tcatgcctct ctcttatgct tttgtggaat
2581 gaggttcctc ggtgtggagg gagggttgga aaacccaaag gaagaaaaag aaatctatgt
2641 tatcccaccc tacctctcac aagcctttcc tgctttaccc ctcacctggc ctctgcccca
2701 cattccttca gcccctcatt tcgagcattg gatttgaggc ttaaggattc aaaaagtcgt
2761 catgaatata gctgatgatt ttatagtggt tctgaaatgg gtcggggatt tgggaacagg
2821 gtggtagtat aagaacaact gatactgttc tctaagctaa atcttagctt ccagctacct
2881 gtcttagatg tggctcttgg gaaccttaga gtgatagcta catagaagtg tgtgggtgtg
2941 tgtgtgtgtg tctgtgtgtg tgtgtgtgag agagagacag acagaaagag agcaagagag
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3061 ctccattaac aggataatcc tcacacctgt ccacatacct gtagtttgtc cttggggatt
3121 ttgaaaattt ttcctccctc tccactccca aactcccaac tcaattaaat gataaaggaa
3181 taggcaaata ggaaaataaa ttagtaaaac ttaagtcaaa gaataggtta ttcatacgct
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3301 ggtacaaggg aggccagaag acgagtggtt cttctctgag gtggacatta aaaaaagaag
3361 aaaatgaagg ggaacctttt gacaagaatg tcaccccaaa ctggattttc atgctgtggt
3421 gtggggaatt ttctgttgtc ctcacttagg tgctggggca gtggtgttag tgatgggtaa
3481 aaaggtagga agctgtcaca gaatcactaa accagggttc ttaacttgtc tgtctataca
3541 tctctgaaat tgggttgaag ttgtgtgcat cattttgagt gacgcactga gaacattcct
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3661 gtcctgctta ctggttttta gtaacaaatg gcagagtatt tctctctgtc tctctctctt
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3841 gcctctttag tagctgggac tacagcatga gccactgccc ttggctaatt tttaaattat
3901 ttttttgtag agatggaaac ttgctatgtt gcccaggcta gtctcaaact cctggactca
3961 agcgatcctc ctaccttggc ctcccaaagt gctgagatta cagtgtgatc cacaccacac
4021 ctggccaaag attggagtat ttttattgct attgttgtgc tgggtgggtg ggtgggtgta

4081 tgctttgtgg ggacgtgtgt tgttgccaag ggctaaatca gttcctaccc tgctgcccac
4141 agtcctccac agctttcctg ctctgtgaag ctaaggatac accccgatga taagctgtca
4201 acata

SEQ ID NO. 5

NM_002964
Homo sapiens S100 calcium binding protein A8 (calgranulin A) S100A8, mRNA

1 atgtctcttg tcagctgtct ttcagaagac ctggtggggc aagtccgtgg gcatcatgtt
61 gaccgagctg gagaaagcct tgaactctat catcgacgtc taccacaagt actccctgat
121 aaaggggaat ttccatgccg tctacaggga tgacctgaag aaattgctag agaccgagtg
181 tcctcagtat atcaggaaaa agggtgcaga cgtctggttc aaagagttgg atatcaacac
241 tgatggtgca gttaacttcc aggagttcct cattctggtg ataaagatgg gcgtggcagc
301 ccacaaaaaa agccatgaag aaagccacaa agagtagctg agttactggg cccagaggct
361 gggcccctgg acatgtacct gcagaataat aaagtcatca atacctcaaa aaaaaaaaaa
421 aaaaaaaa

SEQ ID NO. 6

P06702
Calgranulin B/MRP-14

1 mtckmsqler nietiintfh qysvklghpd tlnqgefkel vrkdlqnflk kenknekvie
61 himedldtna dkqlsfeefi mlmarltwas hekmhegdeg pghhhkpglg egtp

SEQ ID NO. 7

X06233
Human mRNA for calcium-binding protein in macrophages (MRP-14) macrophage migration inhibitory factor (MIF)-related protein

aaaacactct gtgtggctcc tcggctttga cagagtgcaa gacgatgact tgcaaaatgt
cgcagctgga acgcaacata gagaccatca tcaacacctt ccaccaatac tctgtgaagc
tggggcaccc agacaccctg aaccaggggg aattcaaaga gctggtgcga aaagatctgc
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taacctgggc ctcccacgag aagatgcacg agggtgacga gggccctggc caccaccata
agccaggcct cggggagggc acccccctaag accacagtgg ccaagatcac agtggccacg
gccacggcca cagtcatggt ggccacggcc acagccaccc at

SEQ ID NO. 8

M21064
Human migration inhibitory factor-related protein 14 (MRP14) gene, complete cds

1 atcactgtgg agtaggggaa gggcactcct ggggtggcaa ggtgggaggt gggccctgtg
61 ttcccacagt gggcagggag gtagtgaaag ggaagctggc cggacaggaa gggccattcc
121 aagagggctt tgtgcgcagg gctaagccaa gctttctcca taggcaatgg ggagcaactg

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 181 gaggttcgta gcaggagaag gacacatcaa gcccaccagg aggctaagta aaaacagttg
 241 tctcccaagt tataagttcc tggaaccctt gctgggagca ggatttagaa aaatgatgct
 301 gagagatgct agaaacatat tcgccctgag gctctctcac tcagactgca agaggaaggt
 361 atcatcagaa ttgcccttaa ccaggaacca gaatagctgg gtccccttcc tgccaagtca
 421 gcaaccagct atgtgacctt gctcaggtcc atctccgggt gtcagtttct tcatctacaa
 481 tgcaagaggg ttgcccacct ctgagaaccc ttctaacccc aaatctcacc ctatgaatct
 541 aagaacacaa cccctcgcca tcctaagtat cacagagcca ggcaagcatg ggtgagagct
 601 cagaccatcc ttgttggact aaaaggaagg ggcagactgc catgggggc agccgagagg
 661 gtcaggcccc cataggtcct cagcctgctt caacctcaaa ggggatgggg ggctgagtgg
 721 tgccagagga gcagcaggct cgctcgggga gagtagggcc ttaggataga agggaaatga
 781 actaaacaac cagcttcctg caaaccagtt tcaggccagg gctgggaatt tcacaaaaaa
 841 gcagaaggcg ctctgtgaac atttcctgcc ccgccccagc ccccttcctg gcagcattag
 901 cacactgctc acctgtgaag caatcttccg gagacagggc caaagggcaa gtgccccagt
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1501 tgaagctggg gcacccagac accctgaacc agggggaatt caaagagctg gtgcgaaaag
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3121 aagttgttat aaggaccaaa tgaggtgaca tttccaggct tactcatgcc atgaccaggg
3181 caagaccctg gaactcagct tcctcttcta taaatagaga atcagcaccc aagtcacagg
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4021 ccccactgtt cagatcccag gctccctgcc ccatctcaga caccctgtcc agcctgtcca
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4141 cttgatatgc ctgctacctc accaaccagc cccaagcctg tcttcaccca tcactgtcta
4201 cacagccctc tctctctcct aacagaattc tattcctctg aaagtcttca gaaactggac
4261 ctagatagtg ccatgtctgg ggaggaatat ggcaccaggc agtggaaaca aggacagatc
4321 ggtgtgttat ctcacatttg atcagagagc atgatctctc ttaacagacc tgccacccta
4381 atcaacggga gtgctcacac aagtgggagt ctgagagctt agccctatgc ccaccctgg
```

SEQ ID NO. 9

P01833 and Q81ZY7
Polymeric-immunoglobulin receptor (precursor)

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  1 mllfvltcll avfpaistks pifgpeevns vegnsvsitc yypptsvnrh trkywcrqga
 61 rggcitliss egyvsskyag ranltnfpen gtfvvniaql sqddsgrykc glginsrgls
121 fdvslevsqg pgllndtkvy tvdlgrtvti ncpfktenaq krkslykqig lypvlvidss
181 gyvnpnytgr irldiqgtgq llfsvvinql rlsdagqylc qagddsnsnk knadlqvlkp
241 epelvyedlr gsvtfhcalg pevanvakfl crqssgencd vvvntlgkra pafegrilln
301 pqdkdgsfsv vitglrkeda grylcgahsd gqlqegspiq awqlfvnees tiprsptvvk
361 gvagssvavl cpynrkesks ikywclwega qngrcpllvd segwvkaqye grlslleepg
421 ngtftvilnq ltsrdagfyw cltngdtlwr ttveikiieg epnlkvpgnv tavlgetlkv
481 pchfpckfss yekywckwnn tgcqalpsqd egpskafvnc densrlvslt lnlvtradeg
541 wywcgvkqgh fygetaavyv aveerkaags rdvslakada apdekvldsg freienkaiq
601 dprlfaeeka vadtrdqadg srasvdsgss eeqggssral vstlvplglv lavgavavgv
661 ararhrknvd rvsirsyrtd ismsdfensr efgandnmga ssitqetslg gkeefvatte
721 sttetkepkk akrsskeeae maykdfllqs stvaaeaqdg pqea
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SEQ ID NO. 10

NM_002644
Homo sapiens polymeric immunoglobulin receptor (PIGR), mRNA

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   1 agagtttcag ttttggcagc agcgtccagt gccctgccag tagctcctag agaggcaggg
  61 gttaccaact ggccagcagg ctgtgtccct gaagtcagat caacgggaga gaaggaagtg
 121 gctaaaacat tgcacaggag aagtcggcct gagtggtgcg gcgctcggga cccaccagca
 181 atgctgctct tcgtgctcac ctgcctgctg gcggtcttcc cagccatctc cacgaagagt
 241 cccatatttg gtcccgagga ggtgaatagt gtggaaggta actcagtgtc catcacgtgc
 301 tactacccac ccacctctgt caaccggcac acccggaagt actggtgccg gcagggagct
 361 agaggtggct gcataaccct catctcctcg gagggctacg tctccagcaa atatgcaggc
 421 agggctaacc tcaccaactt cccggagaac ggcacatttg tggtgaacat tgcccagctg
 481 agccaggatg actccgggcg ctacaagtgt ggcctgggca tcaatagccg aggcctgtcc
 541 tttgatgtca gcctggaggt cagccagggt cctgggctcc taaatgacac taaagtctac
 601 acagtggacc tgggcagaac ggtgaccatc aactgccctt tcaagactga gaatgctcaa
 661 aagaggaagt ccttgtacaa gcagataggc ctgtaccctg tgctggtcat cgactccagt
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 721 ggttatgtaa atcccaacta tacaggaaga atacgccttg atattcaggg tactggccag
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 841 caggctgggg atgattccaa tagtaataag aagaatgctg acctccaagt gctaaagccc
 901 gagcccgagc tggtttatga agacctgagg ggctcagtga ccttccactg tgccctgggc
 961 cctgaggtgg caaacgtggc caaatttctg tgccgacaga gcagtgggga aaactgtgac
1021 gtggtcgtca acaccctggg gaagagggcc ccagcctttg agggcaggat cctgctcaac
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1321 atcaagtact ggtgtctctg ggaaggggcc cagaatggcc gctgcccccc gctggtggac
1381 agcgaggggt gggttaaggc ccagtacgag ggccgcctct ccctgctgga ggagccaggc
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1501 tgtctgacca acggcgatac tctctggagg accaccgtgg agatcaagat tatcgaagga
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1921 gctcctgatg agaaggtgct agactctggt tttcgggaga ttgagaacaa agccattcag
1981 gatcccaggc tttttgcaga ggaaaaggcg gtggcagata caagagatca agccgatggg
2041 agcagagcat ctgtggattc cggcagctct gaggaacaag gtggaagctc cagagcgctg
2101 gtctccaccc tggtgccccc gggcctggtg ctggcagtgg gagccgtggc tgtgggggtg
2161 gccagagccc ggcacaggaa gaacgtcgac cgagtttcaa tcagaagcta caggacagac
2221 attagcatgt cagacttcga gaactccagg gaatttggag ccaatgacaa catgggagcc
2281 tcttcgatca ctcaggagac atccctcgga ggaaaagaag agtttgttgc caccactgag
2341 agcaccacag agaccaaaga acccaagaag gcaaaaaggt catccaagga ggaagccgag
2401 atggcctaca aagacttcct gctccagtcc agcaccgtgg ccgccgaggc ccaggacggc
2461 ccccaggaag cctagacggt gtcgccgcct gctccctgca cccatgacaa tcaccttcag
2521 aatcatgtcg atcctggggc cctcagctcc tggggacccc actccctgct ctaacacctg
2581 cctaggtttt tcctactgtc ctcagaggcg tgctggtccc ctcctcagtg acatcaaagc
2641 ctggcctaat tgttcctatt ggggatgagg gtggcatgag gaggtcccac ttgcaacttc
2701 tttctgttga gagaacctca ggtacggaga agaatagagg tcctcatggg tcccttgaag
2761 gaagagggac cagggtggga gagctgattg cagaaaggag agacgtgcag cgcccctctg
2821 cacccttatc atgggatgtc aacagaattt ttccctccac tccatccctc cctcccgtcc
2881 ttcccctctt cttctttcct tccatcaaaa gatgtatttg aattcatact agaattcagg
2941 tgctttgcta gatgctgtga caggtatgcc accaacactg ctcacagcct ttctgaggac
3001 accagtgaaa gaagccacag ctcttcttgg cgtatttata ctcactgagt cttaactttt
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3241 ttttttgaga cggagtctct cactgtcgcc caggctggag tgcagtggcg caatctcggc
3301 tcactgcaac ctccgcctcc caggttgaca ccattctcct gcctcaccct cccaagtagc
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3481 gcctcccaaa gtgctgggat tacaggcgtg agccaccgcg tccggcctct ttttttcttt
3541 tctttttttt gagacaaagt ctcactgtgt cacccagact ggaatgcagt gacacaatct
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3661 tagctgggac tacagatgtg ggccaccatg tctggctaat tttttttttt tttttttttt
3721 tttgtagaga cagggtttcg ccatgttgac gagactggtc tcgaactcct ggcctcaagt
3781 gatctgccgc ctcagcttct caaagtactg ggattatata ggcatgagcc actgagcctg
3841 gccctgaagc gtttttctca aaggccctca gtgagataaa ttagatttgg catctcctgt
3901 cctgggccag ggatctctct acaagagccc ctgcccctct gttggaggca cagttttaga
3961 ataaggagga ggagggagaa gagaaaatgt aaaggaggga gatctttccc aggccgcacc
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4021 atttctgtca ctcacatgga cccaagataa aagaatggcc aaaccctcac aacccctgat
4081 gtttgaagag ttccaagttg aagggaaaca aagaagtgtt tgatggtgcc agagaggggc
4141 tgctctccag aaagctaaaa tttaatttct tttttcctct gagttctgta cttcaaccag
4201 cctacaagct ggcacttgct aacaaatcag aaatatgaca attaatgatt aaagactgtg
4261 attgcc

SEQ ID NO. 11

P30086 - Homo sapiens
Phosphatidylethanolamine binding protein (PEBP)

1 mpvdlskwsg plslqevdeq pqhplhvtya gaavdelgkv ltptqvknrp tsiswdglds
61 gklytlvltd pdapsrkdpk yrewhhflvv nmkgndissg tvlsdyvgsg ppkgtglhry
121 vwlvyeqdrp lkcdepilsn rsgdhrgkfk vasfrkkyel rapvagtcyq aewddyvpkl
181 yeqlsgk

SEQ ID NO. 12

NM_002567
Homo sapiens prostatic binding protein (PBP), mRNA

1 tgggcggcgg ctgaggcgcg tgctctcgcg tggtcgctgg gtctgcgtct tcccgagcca
61 gtgtgctgag ctctccgcgt cgcctctgtc gcccgcgcct ggcctaccgc ggcactcccg
121 gctgcacgct ctgcttggcc tcgccatgcc ggtggacctc agcaagtggt ccgggccctt
181 gagcctgcaa gaagtggacg agcagccgca gcacccgctg catgtcacct acgccggggc
241 ggcggtggac gagctgggca aagtgctgac gcccacccag gttaagaata gacccaccag
301 catttcgtgg gatggtcttg attcagggaa gctctacacc ttggtcctga cagacccgga
361 tgctcccagc aggaaggatc ccaaatacag agaatggcat catttcctgg tggtcaacat
421 gaagggcaat gacatcagca gtggcacagt cctctccgat tatgtgggct cggggcctcc
481 caagggcaca ggcctccacc gctatgtctg gctggtttac gagcaggaca ggccgctaaa
541 gtgtgacgag cccatcctca gcaaccgatc tggagaccac cgtggcaaat tcaaggtggc
601 gtccttccgt aaaaagtatg agctcagggc cccggtggct ggcacgtgtt accaggccga
661 gtgggatgac tatgtgccca aactgtacga gcagctgtct gggaagtagg gggttagctt
721 ggggacctga actgtcctgg aggccccaag ccatgttccc cagttcagtg ttgcatgtat
781 aatagatttc tcctcttcct gccccccttg gcatgggtga gacctgacca gtcagatggt
841 agttgagggt gacttttcct gctgcctggc ctttataatt ttactcactc actctgattt
901 atgttttgat caaatttgaa cttcattttg gggggtattt tggtactgtg atggggtcat
961 caaattatta atctgaaaat agcaacccag aatgtaaaaa agaaaaaact gggggggaaaa
1021 agaccaggtc tacagtgata gagcaaagca tcaaagaatc tttaagggag gtttaaaaaa
1081 aaaaaaaaaa aaaaagattg gttgcctctg cctttgtgat cctgagtcca gaatggtaca
1141 caatgtgatt ttatggtgat gtcactcacc tagacaacca gaggctggca ttgaggctaa
1201 cctccaacac agtgcatctc agatgcctca gtaggcatca gtatgtcact ctggtccctt
1261 taaagagcaa tcctggaaga agcaggaggg agggtggctt tgctgttgtt gggacatggc
1321 aatctagacc ggtagcagcg ctcgctgaca gcttgggagg aaacctgaga tctgtgtttt
1381 ttaaattgat cgttcttcat gggggtaaga aaagctggtc tggagttgct gaatgttgca
1441 ttaattgtgc tgtttgcttg tagttgaata aaaatagaaa cctgaatgaa gaaaaaaaaa
1501 aaaaaaa

SEQ ID NO. 13

P39687 - Homo sapiens
Acidic leucine-rich nuclear phosphoprotein 32 family member A

```
  1 memgrrihle lrnrtpsdvk elvldnsrsn egklegltde feeleflsti nvgltsianl
 61 pklnklkkle lsdnrvsggl evlaekcpnl thlnlsgnki kdlstieplk klenlksldl
121 fncevtnlnd yrenvfkllp qltyldgydr ddkeapdsda egyvegldde eededeeeyd
181 edaqvvedee dedeeeegee edvsgeeeed eegyndgevd deedeeelge eergqkrkre
241 pedegeddd
```

SEQ ID NO. 14

NM_006305
Homo sapiens acidic (leucine-rich) nuclear phosphoprotein 32 family, member A (ANP32A), mRNA

```
  1 cgggtgctgg gggctcgaga accgagcgga gctggttgag ccttcaaagt cctaaaacgc
 61 gcggccgtgg gttcggggtt tattgattga attccgccgg cgcgggagcc tctgcagaga
121 gagagcgcga gagatggaga tgggcagacg gattcattta gagctgcgga acaggacgcc
181 ctctgatgtg aaagaacttg tcctggacaa cagtcggtcg aatgaaggca aactcgaagg
241 cctcacagat gaatttgaag aactggaatt cttaagtaca atcaacgtag gcctcacctc
301 aatcgcaaac ttaccaaagt taaacaaact taagaagctt gaactaagcg ataacagagt
361 ctcaggggge ctggaagtat tggcagaaaa gtgtccgaac ctcacgcatc taaatttaag
421 tggcaacaaa attaaagacc tcagcacaat agagccactg aaaaagttag aaaacctcaa
481 gagcttagac cttttcaatt gcgaggtaac caacctgaac gactaccgag aaaatgtgtt
541 caagctcctc ccgcaactca catatctcga cggctatgac cgggacgaca aggaggcccc
601 tgactcggat gctgagggct acgtggaggg cctggatgat gaggaggagg atgaggatga
661 ggaggagtat gatgaagatg ctcaggtagt ggaagacgag gaggacgagg atgaggagga
721 ggaaggtgaa gaggaggacg tgagtggaga ggaggaggag gatgaagaag gttataacga
781 tggagaggta gatgacgagg aagatgaaga agagcttggt gaagaagaaa ggggtcagaa
841 gcgaaaacga gaacctgaag atgagggaga agatgatgac taagtggaat aacctatttt
901 gaaaaattcc tattgtgatt tgactgtttt tacccatatc ccctctcccc ccccctcca
961 atcctgcccc ctgaaactta ttttttttctg attgtaacgt tgctgtggga acgagagggg
1021 aagagtgtac tgggggttgc gggggggaggg atggcgggtg ggggtggaat aaaatactat
1081 ttttactgcc actctttaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaa
```

SEQ ID NO. 15

P17066 - Homo sapiens
Heat shock 70kDa protein

```
  1 mqaprelavg idlgttyscv gvfqqgrvei landqgnrtt psyvaftdte rlvgdaaksq
 61 aalnphntvf dakrligrkf adttvqsdmk hwpfrvvseg gkpkvrvcyr gedktfypee
121 issmvlskmk etaeaylgqp vkhavitvpa yfndsqrqat kdagaiagln vlriinepta
181 aaiaygldrr gagernvlif dlgggtfdvs vlsidagvfe vkatagdthl ggedfdnrlv
241 nhfmeefrrk hgkdlsgnkr alrrlrtace rakrtlssst qatleidslf egvdfytsit
301 rarfeelcsd lfrstlepve kalrdakldk aqihdvvlvg gstripkvqk llqdffngke
361 lnksinpdea vaygaavqaa vlmgdkcekv qdllldvap lslgletagg vmttliqrna
421 tiptkqtqtf ttysdnqpgv fiqvyegera mtkdnnllgr felsgippap rgvpqievtf
481 didangilsv tatdrstgka nkititndkg rlskeeverm vheaeqykae deaqrdrvaa
541 knsleahvfh vkgslqeesl rdkipeedrr kmqdkcrevl awlehnqlae keeyehqkre
601 leqicrpifs rlyggpgvpg gsscgtqarq gdpstgpiie evd
```

SEQ ID NO. 16

NM_002155
Homo sapiens heat shock 70kDa protein 6 (HSP70B') (HSPA6), mRNA.

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   1 agagccagcc cggaggagct agaaccttcc ccgcatttct ttcagcagcc tgagtcagag
  61 gcgggctggc ctggcgtagc cgcccagcct cgcggctcat gccccgatct gcccgaacct
 121 tctcccgggg tcagcgccgc gccgcgccac ccggctgagt cagcccgggc gggcgagagg
 181 ctctcaactg ggcgggaagg tgcgggaagg tgcggaaagg ttcgcgaaag ttcgcggcgg
 241 cgggggtcgg gtgaggcgca aaaggataaa aagcccgtgg aagcggagct gagcagatcc
 301 gagccgggct ggctgcagag aaaccgcagg gagagcctca ctgctgagcg cccctcgacg
 361 gcggagcggc agcagcctcc gtggcctcca gcatccgaca agaagcttca gccatgcagg
 421 ccccacggga gctcgcggtg ggcatcgacc tgggcaccac ctactcgtgc gtgggcgtgt
 481 ttcagcaggg ccgcgtggag atcctggcca acgaccaggg caaccgcacc acgcccagct
 541 acgtggcctt caccgacacc gagcggctgg tcggggacgc ggccaagagc caggcggccc
 601 tgaaccccca caacaccgtg ttcgatgcca agcggctgat cgggcgcaag ttcgcggaca
 661 ccacggtgca gtcggacatg aagcactggc ccttccgggt ggtgagcgag ggcggcaagc
 721 ccaaggtgcg cgtatgctac cgcggggagg acaagacgtt ctaccccgag gagatctcgt
 781 ccatggtgct gagcaagatg aaggagacgg ccgaggcgta cctgggccag cccgtgaagc
 841 acgcagtgat caccgtgccc gcctatttca atgactcgca gcgccaggcc accaaggacg
 901 cgggggccat cgcggggctc aacgtgttgc ggatcatcaa tgagcccacg gcagctgcca
 961 tcgcctatgg gctggaccgg cggggcgcgg gagagcgcaa cgtgctcatt tttgacctgg
1021 gtgggggcac cttcgatgtg tcggttctct ccattgacgc tggtgtcttt gaggtgaaag
1081 ccactgctgg agatacccac ctgggaggag aggacttcga caaccggctc gtgaaccact
1141 tcatggaaga attccggcgg aagcatggga aggacctgag cgggaacaag cgtgccctgc
1201 gcaggctgcg cacagcctgt gagcgcgcca agcgcaccct gtcctccagc acccaggcca
1261 ccctggagat agactccctg ttcgagggcg tggacttcta cacgtccatc actcgtgccc
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1381 tgcgggatgc caagctggac aaggcccaga ttcatgacgt cgtcctggtg gggggctcca
1441 ctcgcatccc caaggtgcag aagttgctgc aggacttctt caacggcaag gagctgaaca
1501 agagcatcaa ccctgatgag gctgtggcct atggggctgc tgtgcaggcg gccgtgttga
1561 tgggggacaa atgtgagaaa gtgcaggatc tcctgctgct ggatgtggct cccctgtctc
1621 tggggctgga gacagcaggt ggggtgatga ccacgctgat ccagaggaac gccactatcc
1681 ccaccaagca gacccagact ttcaccacct actcggacaa ccagcctggg gtcttcatcc
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1861 atgctaatgg catcctgagc gtgacagcca ctgacaggag cacaggtaag gctaacaaga
1921 tcaccatcac caatgacaag ggccggctga gcaaggagga ggtggagagg atggttcatg
1981 aagccgagca gtacaaggct gaggatgagg cccagaggga cagagtggct gccaaaaact
2041 cgctggaggc ccatgtcttc catgtgaaag gttctttgca agaggaaagc cttagggaca
2101 agattcccga agaggacagg cgcaaaatgc aagacaagtg tcgggaagtc cttgcctggc
2161 tggagcacaa ccagctggca gagaaggagg agtatgagca tcagaagagg gagctggagc
2221 aaatctgtcg ccccatcttc tccaggctct atgggggcc tggtgtccct gggggcagca
2281 gttgtggcac tcaagcccgc caggggggacc ccagcaccgg ccccatcatt gaggaggttg
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2461 cttcccagga taactgaagt cttttgactt tttgggggga gggcggttca tcctcttctg
2521 cttcaaataa aaagtcatta atttattaaa acttgtgtgg cactttaaca ttgctttcac
2581 ctatattttg tgtactttgt tacttgcatg tatgaatttt gttatgtaaa atatagttat
2641 agacctaaat aaaaaaaaaa aaaa
```

SEQ ID NO. 17

X51757
Human heat-shock protein HSP70B gene

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   1 cccgggcggg cgagaggctc tcaactgggc gggaaggtgc gggaaggtgc ggaaaggttc
  61 gcgaaagttc gcggcggcgg gggtcgggtg aggcgcaaaa ggataaaaag cccgtggaag
 121 cggagctgag cagatccgag ccgggctggc tgcagagaca ccgcagggag agcctcactg
 181 ctgagcgccc ctcgacggcg gacgggcagc agcctccgtg gcctccagca tccgacaaga
 241 agcttcagcc atgcaggccc cacgggagct cgcggtgggc atcgacctgg gcaccaccta
 301 ctcgtgcgtg ggcgtgtttc agcagggccg cgtggagatc ctggccaacg accagggcaa
 361 ccgcaccacg cccagctacg tggccttcac cgacaccgag cggctggtcg gggacgcggc
 421 caagagccag gcggccctga acccccacaa caccgtgttc gatgccaagc ggctgatcgg
 481 gcgcaagttc gcggacacca cggtgcagtc ggacatgaag cactggccct tccgggtggt
 541 gagcgagggc ggcaagccca aggtgccggt atcgtaccgc ggggaggaca agacgttcta
 601 ccccgaggag atctcgtcca tggtgctgag caagatgaag gagacggccg aggcgtacct
 661 gggccagccc gtgaagcacg cagtgatcac cgtgcccgcc tatttcaatg actcgcagcg
 721 ccaggccacc aaggacgcgg gggccatcgc ggggctcaac gtgttgcgga tcatcaatga
 781 gcccacggca gctgccatcg cctatgggct ggaccggcgg ggcgcgggag agcgcaacgt
 841 gctcattttt gacctgggtg ggggcacctt cgatgtgtcg gttctctcca ttgacgctgg
 901 tgtctttgag gtgaaagcca ctgctggaga tacccacctg ggaggagagg acttcgacaa
 961 ccggctcgtg aaccacttca tggaagaatt ccggcggaag catgggaagg acctgagcgg
1021 gaacaagcgt gccctcggca ggctgcgcac agcctgtgag cgcgccaagc gcaccctgtc
1081 ctccagcacc caggccaccc tggagataga ctccctgttc gagggcgtgg acttctacac
1141 gtccatcact cgtgcccgct ttgaggaact gtgctcagac ctcttccgca gcaccctgga
1201 gccggtggag aaggccctgc gggatgccaa gctggacaag gcccagattc atgacgtcgt
1261 cctggtgggg ggctccactc gcatccccaa ggtgcagaag ttgctgcagg acttcttcaa
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1561 gcctggggtc ttcatccagg tgtatgaggg tgagagggcc atgaccaagg acaacaacct
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1681 ggtgaccttt gacattgatg ctaatggcat cctgagcgtg acagccactg acaggagcac
1741 aggtaaggct aacaagatca ccatcaccaa tgacaagggc cggctgagca aggaggaggt
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1921 ggaaagcctt agggacaaga ttcccgaaga ggacaggcgc aaaatgcaag acaagtgtcg
1981 ggaagtcctt gcctggctgg agcacaacca gctggcagag aaggaggagt atgagcatca
2041 gaagagggag ctggagcaaa tctgtcgccc catcttctcc aggctctatg gggggcctgg
2101 tgtccctggg ggcagcagtt gtggcactca agcccgccag ggggacccca gcaccggccc
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2221 tatgctatgg gccttctaga ctgtcttcta tgatcctgcc cttcagagat gaactttccc
2281 tccaaagcta gaactttctt cccaggataa ctgaagtctt ttgacttttt gcggggaggg
2341 cggttcatcc tcttctgctt caaataaaaa gtcattaatt tattaaaact tgtgtggcac
2401 tttaacattg ctttcaccta tattttgtgt actttgttac ttgcatgtat gaattttgtt
2461 atgtaaaata tagttataga cctaaataag ct
```

SEQ ID NO. 18

P14174
macrophage migration inhibitory factor - Homo Sapiens

```
 1 mpmfivntnv prasvpdgfl seltqqlaqa tgkppqyiav hvvpdqlmaf ggssepcalc
61 slhsigkigg aqnrsyskll cgllaerlri spdrvyinyy dmnaanvgwn nstfa
```

SEQ ID NO. 19

NM_002415 - Homo Sapiens
Homo sapiens macrophage migration inhibitory factor (glycosylation-inhibiting factor) (MIF), mRNA

```
   1 accacagtgg tgtccgagaa gtcaggcacg tagctcagcg gcggccgcgg cgcgtgcgtc
  61 tgtgcctctg cgcgggtctc ctggtccttc tgccatcatg ccgatgttca tcgtaaacac
 121 caacgtgccc cgcgcctccg tgccggacgg gttcctctcc gagctcaccc agcagctggc
 181 gcaggccacc ggcaagcccc cccagtacat cgcggtgcac gtggtcccgg accagctcat
 241 ggccttcggc ggctccagcg agccgtgcgc gctctgcagc ctgcacagca tcggcaagat
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 361 gcgcatcagc ccggacaggg tctacatcaa ctattacgac atgaacgcgg ccaatgtggg
 421 ctggaacaac tccaccttcg cctaagagcc gcagggaccc acgctgtctg cgctggctcc
 481 acccgggaac ccgccgcacg ctgtgttcta ggcccgccca ccccaacctt ctggtgggga
 541 gaaataaacg gtttagagac t
```

SEQ ID NO. 20

L19686
Homo sapiens macrophage migration inhibitory factor (MIF) gene, complete cds

```
   1 ctgcaggaac caatacccat aggctatttg tataaatggg ccatggggcc tcccagctgg
  61 aggctggctg gtgccacgag ggtcccacag gcatgggtgt ccttcctata tcacatggcc
 121 ttcactgaga ctggtatatg gattgcacct atcagagacc aaggacagga cctccctgga
 181 aatctctgag gacctggcct gtgatccagt tgctgccttg tcctcttcct gctatgtcat
 241 ggcttatctt ctttcaccca ttcattcatt cattcattca ttcagcagta ttagtcaatg
 301 tctcttgata tgcctggcac ctgctagatg gtccccgagt ttaccattag tggaaaagac
 361 atttaagaaa ttcaccaagg gctctatgag aggccataca cggtggacct gactagggtg
 421 tggcttccct gaggagctga agttgcccag aggcccagag aaggggagct gagcacgttt
 481 gaaccactga acctgctctg gacctcgcct ccttccttcg gtgcctccca gcatcctatc
 541 ctctttaaag agcaggggtt cagggaagtt ccctggatgg tgattcgcag gggcagctcc
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 781 cgcatcccca cccccggcac agcgcctcct ggcgactaac atcggtgact tagtgaaagg
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 961 gggtcctgcc ctgcgggggt cgagccgagg caggcggtga cttccccact cggggcggag
1021 ccgcagcctc gcgggggcgg ggcctggcgc cggcggtggc gtcacaaaag gcgggaccac
1081 agtggtgtcc gagaagtcag gcacgtagct cagcggcggc cgcggcgcgt gcgtctgtgc
1141 ctctgcgcgg gtctcctggt ccttctgcca tcatgccgat gttcatcgta aacaccaacg
1201 tgccccgcgc ctccgtgccg gacgggttcc tctccgagct cacccagcag ctggcgcagg
1261 ccaccggcaa gccccccccag gtttgccggg aggggacagg aagaggggg tgcccaccgg
1321 acgaggggtt ccgcgctggg agctggggag gcgactcctg aacggagctg gggggcgggg
1381 cgggggggagg acggtggctc gggcccgaag tggacgttcg gggcccgacg aggtcgctgg
1441 ggcgggctga ccgcgccctt tcctcgcagt acatcgcggt gcacgtggtc ccggaccagc
1501 tcatggcctt cggcggctcc agcgagccgt gcgcgctctg cagcctgcac agcatcggca
1561 agatcggcgg cgcgcagaac cgctcctaca gcaagctgct gtgcggcctg ctggccgagc
1621 gcctgcgcat cagcccggac aggtacgcgg agtcgcggag gggcgggggga ggggcggcgg
1681 cgcgcggcca ggcccgggac tgagccaccc gctgagtccg gcctcctccc cccgcagggt
1741 ctacatcaac tattacgaca tgaacgcggc caatgtgggc tggaacaact ccaccttcgc
1801 ctaagagccg cagggaccca cgctgtctgc gctggctcca cccgggaacc cgccgcacgc
1861 tgtgttctag gcccgcccac cccaaccttc tggtggggag aaataaacgg tttagagact
1921 aggagtgcct cggggttcct tggcttgcgg gaggaattgg tgcagagccg ggacattggg
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1981 gagcgaggtc gggaaacggt gttgggggcg ggggtcaggg ccgggttgct ctcctcgaac
2041 ctgctgttcg ggagcccttt tgtccagcct gtccctccta cgctcctaac agaggagccc
2101 cagtgtcttt ccattctatg gcgtacgaag ggatgaggag aagttggcac tctgccctgg
2161 gctgcag

SEQ ID NO. 21

P31949
Calgizzarin - Homo sapiens

1 makissptet erciesliav fqkyagkdgy nytlsktefl sfmntelaaf tknqkdpgvl
61 drmmkkldtn sdgqldfsef lnligglama chdsflkavp sqkrt

SEQ ID NO. 22

NM_005620 and D38583 - Homo sapiens
Homo sapiens S100 calcium binding protein A11 (calgizzarin)(S100A11), mRNA

1 gggcaaggct gggccgggaa gggcgtgggt tgaggagagg ctccagaccc gcacgccgcg
61 cgcacagagc tctcagcgcc gctcccagcc acagcctccc gcgcctcgct cagctccaac
121 atggcaaaaa tctccagccc tacagagact gagcggtgca tcgagtccct gattgctgtc
181 ttccagaagt atgctggaaa ggatggttat aactacactc tctccaagac agagttccta
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301 gaccgcatga tgaagaaact ggacaccaac agtgatggtc agctagattt ctcagaattt
361 cttaatctga ttggtggcct agctatggct tgccatgact ccttcctcaa ggctgtccct
421 tcccagaagc ggacctgagg acccccttggc cctggccttc aaacccaccc cctttccttc
481 cagcctttct gtcatcatct ccacagccca cccatccccc gagcacacta accacctcat
541 gcaggcccca cctgccaata gtaataaagc aatgtcactt ttttaaaaca tgaaa

SEQ ID NO. 23

P00938 and NP_000356 - Homo sapiens
Triosephosphate isomerase

1 mapsrkffvg gnwkmngrkq slgeligtln aakvpadtev vcapptayid farqkldpki
61 avaaqncykv tngaftgeis pgmikdcgat wvvlghserr hvfgesdeli gqkvahalae
121 glgviacige kldereagit ekvvfeqtkv iadnvkdwsk vvlayepvwa igtgktatpq
181 qaqevheklr gwlksnvsda vaqstriiyg gsvtgatcke lasqpdvdgf lvggaslkpe
241 fvdiinakq

SEQ ID NO. 24

NM_000365
Homo sapiens triosephosphate isomerase 1 (TPI1), mRNA

1 ccttcagcgc ctcggctcca gcgccatggc gccctccagg aagttcttcg ttgggggaaa
61 ctggaagatg aacgggcgga agcagagtct gggggagctc atcggcactc tgaacgcggc
121 caaggtgccg gccgacaccg aggtggtttg tgctcccccc actgcctata tcgacttcgc
181 ccggcagaag ctagatccca agattgctgt ggctgcgcag aactgctaca aagtgactaa
241 tggggctttt actggggaga tcagccctgg catgatcaaa gactgcggag ccacgtgggt

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 301 ggtcctgggg cactcagaga gaaggcatgt ctttggggag tcagatgagc tgattgggca
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 421 agatgaaagg gaagctggca tcactgagaa ggttgttttc gagcagacaa aggtcatcgc
 481 agataacgtg aaggactgga gcaaggtcgt cctggcctat gagcctgtgt gggccattgg
 541 tactggcaag actgcaacac cccaacaggc ccaggaagta cacgagaagc tccgaggatg
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 661 tgtgactggg gcaacctgca aggagctggc cagccagcct gatgtggatg gcttccttgt
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 781 atccatcttc cctacccttc ctgccaagcc agggactaag cagcccagaa gcccagtaac
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 961 ccttctccac ttactataat ggttggaact aaacgtcacc aaggtggctt ctccttggct
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1141 tgctgccctc tcccatggtg cccgtgcctc tgtgctgtgt atgtgaacca cccatgtgag
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SEQ ID NO. 25

X69723
H.sapiens TPI1 gene for triosephosphate isomerase.

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   1 ctgcagttcc tgccaggcct tgccagccgg ggcgagggtt gggatgatcc tggcggccta
  61 tgcctgtgtg ggctgcccct cccgctgtga accctgcatt tgtcccgcaa gttttcactc
 121 aggtagactc cctgggtaca agggtgcctg ctcagcagtc gggcatgagc tgctccgatg
 181 ggcgaaggag gttgtctatt ccacagttgg agaggggccc tctctgcccc agtgggcgat
 241 ctgggctacg gccaagttgc caccagctag ttccgcttga aaaccacttc tggccccgtg
 301 ggggactcaa gtcgccaagc gagggttccc ctgagcgccg gagctcacag gtctcgcctt
 361 gtcccgaaag ccccgcaatc gaggcggagg cgaccgagcc cccgactctc ctagaacgtt
 421 gccacaagaa gggggaacgt cggaacagtg catcatcggg cggcggccgg ggcggcggca
 481 ggagggcggg cgggggcagg ggctccgggg gactgggcgg gccatggcgg aggacggcga
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 781 gccggggccg ggccggggca ggagtggcag cgcctctccc gaggcccgag gtccgggccg
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1921 gtttgtgctc cccctactgc ctatatcgac ttcgcccggc agaagctaga tcccaagatt
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2461 acctttgagc ctatccaggg ccacagagac tcagagggta gggtcaggcc ctggagcctg
2521 tcttggtccc catgctgatc cagaaaagga aaaaggggag ggggagtgac aatctttgct
2581 tggggcctat gacttctcca gccccaaggt agatgccacc tggaaatccc ccaatgtcca
2641 ctaggggcca gtaggccacc gttcttcgta ctccggagaa cctggctgga gagctctttc
2701 ttgttcaccc ttccctccat ctgtatctct gccctgcaga taacgtgaag gactggagca
2761 aggtcgtcct ggcctatgag cctgtgtggg ccattggtac tggcaagact gcaacacccc
2821 aacaggtaac cgggcccagg agccctgccc tcatcccagc ctgcctcaat aggtttggac
2881 agacacagcc cacatggagc aaccccttat ttcaaagaca cagagacctt gaacccagag
2941 acagtgactt gtccaagggc atccagtcca gggcctggct tggatcagag ccctggtact
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3661 aaggtggctt ctccttggct gagagatgga aggcgtggtg ggatttgctc ctgggttccc
3721 taggccctag tgagggcaga agagaaacca tcctctccct tcttacaccg tgaggccaag
3781 atcccctcag aaggcaggag tgctgccctc tcccatggtg cccgtgcctc tgtgctgtgt
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3901 cactggactt gcccagataa tcttcctttt tgaggcagct atataaatga tcatttgtgc
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4561 agagctggct aaggccctct aaacaacagg ccaaggtggc tctgacagtg gtggagctgg
4621 cccaggcttt gactccagag gcttgggagc tggggctgag gtgaggaggg atggccctcc
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4861 atgtggcaca gagacaggtt agagcccagg gaatccggta tacagcctgg gtacctcgtc
4921 tgcccatcct tcttttggac ctgtacatca aacccagtac ctaaccgttt gcacctcttg
4981 cctaggggtg attactcctg aattc

SEQ ID NO. 26

Q05586 - Homo sapiens
Glutamate [NMDA] receptor subunit zeta 1 precursor

MSTMRLLTLA LLFSCSVARA ACDPKIVNIG AVLSTRKHEQ MFREAVNQAN KRHGSWKIQL
NATSVTHKPN AIQMALSVCE DLISSQVYAI LVSHPPTPND HFTPTPVSYT AGFYRIPVLG
LTTRMSIYSD KSIHLSFLRT VPPYSHQSSV WFEMMRVYSW NHIILLVSDD HEGRAAQKRL
ETLLEERESK AEKVLQFDPG TKNVTALLME AKELEARVII LSASEDDAAT VYRAAAMLNM
TGSGYVWLVG EREISGNALR YAPDGILGLQ LINGKNESAH ISDAVGVVAQ AVHELLEKEN
ITDPPRGCVG NTNIWKTGPL FKRVLMSSKY ADGVTGRVEF NEDGDRKFAN YSIMNLQNRK
LVQVGIYNGT HVIPNDRKII WPGGETEKPR GYQMSTRLKI VTIHQEPFVY VKPTLSDGTC
KEEFTVNGDP VKKVICTGPN DTSPGSPRHT VPQCCYGFCI DLLIKLARTM NFTYEVHLVA
DGKFGTQERV NNSNKKEWNG MMGELLSGQA DMIVAPLTIN NERAQYIEFS KPFKYQGLTI
LVKKEIPRST LDSFMQPFQS TLWLLVGLSV HVVAVMLYLL DRFSPFGRFK VNSEEEEEDA
LTLSSAMWFS WGVLLNSGIG EGAPRSFSAR ILGMVWAGFA MIIVASYTAN LAAFLVLDRP
EERITGINDP RLRNPSDKFI YATVKQSSVD IYFRRQVELS TMYRHMEKHN YESAAEAIQA
VRDNKLHAFI WDSAVLEFEA SQKCDLVTTG ELFFRSGFGI GMRKDSPWKQ NVSLSILKSH
ENGFMEDLDK TWVRYQECDS RSNAPATLTF ENMAGVFMLV AGGIVAGIFL IFIEIAYKRH
KDARRKQMQL AFAAVNVWRK NLQDRKSGRA EPDPKKKATF RAITSTLASS FKRRRSSKDT
STGGGRGALQ NQKDTVLPRR AIEREEGQLQ LCSRHRES

SEQ ID NO. 27

D13515
Homo sapiens mRNA for key subunit of N-methyl-D-aspartate receptor, complete cds

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   1 gcttcagcgc cccttcctc ggccgacgtc ccgggaccgc cgctccgggg gagacgtggc
  61 gtccgcagcc cgcggggccg ggcgagcgca ggacggcccg gaagccccgc gggggatgcg
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SEQ ID NO. 28

LLTLLALLFSCSVAR

SEQ ID NO. 29

ITMLCTGSRTLK

SEQ ID NO. 30

ITHU and P01009 - Homo sapiens
α-1-antitrypsin precursor

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SEQ ID NO. 31

NM_000295

Homo sapiens serine (or cysteine) proteinase inhibitor, clade A(alpha-1 antiproteinase, antitrypsin), member 1 (SERPINA1),transcript variant 1, mRNA

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SEQ ID NO. 32

K02212
Human alpha-1-antitrypsin gene (S variant), complete cds

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11761 ggagcaaggc ctatgtgaca gggagggaga ggatgtgcag ggccagggcc gtccaggggg
11821 agtgagcgct tcctgggagg tgtccacgtg agccttgctc gaggcctggg atcagcctta
11881 caacgtgtct ctgcttctct cccctccagg ccgtgcataa ggctgtgctg accatcgacg
11941 agaaagggac tgaagctgct ggggccatgt ttttagaggc catacccatg tctatccccc
12001 ccgaggtcaa gttcaacaaa ccctttgtct tcttaatgat tgaacaaaat accaagtctc
12061 ccctcttcat gggaaaagtg gtgaatccca cccaaaaata actgcctctc gctcctcaac
12121 ccctcccctc catccctggc cccctccctg gatgacatta aagaagggtt gagctggtcc
12181 ctgcctgcat gtgatctgta aatccctggg atgttttctc tg
```

SEQ ID NO. 33

gi/125294, P12277 - Homo sapiens
Creatine kinase, B chain (B-CK)

```
mpfsnshnal klrfpaedef pdlsahnnhm akvltpelya elrakstpsg ftlddviqtg
vdnpghpyim tvgcvagdee syevfkdlfd piiedrhggy kpsdehktdl npdnlqggdd
ldpnyvlssr vrtgrsirgf clpphcsrge rraieklave alssldgdla gryyalksmt
eaeqqqlidd hflfdkpvsp lllasgmard wpdargiwhn dnktflvwvn eedhlrvism
qkggnmkevf trfctgltqi etlfkskdye fmwnphlgyi ltcpsnlgtg lragvhiklp
nlgkhekfse vlkrlrlqkr gtggvdtaav ggvfdvsnad rlgfsevelv qmvvdgvkll
iemeqrleqg qaiddlmpaq k
```

SEQ ID NO. 34

NM_001823 Homo sapiens creatine kinase, brain (CKB), mRNA
Creatine kinase, B chain (B-CK)

```
   1 gctgttcgcc tgcgtcgctc cgggagctgc cgacggacgg agcgcccccg ccccgcccg
  61 gccgcccgcc cgccgccgcc atgcccttct ccaacagcca caacgcactg aagctgcgct
 121 tcccggccga ggacgagttc cccgacctga gcgcccacaa caaccacatg gccaaggtgc
 181 tgacccccga gctgtacgcg gagctgcgcg ccaagagcac gccgagcggc ttcacgctgg
 241 acgacgtcat ccagacaggc gtggacaacc cgggccaccc gtacatcatg accgtgggct
 301 gcgtggcggg cgacgaggag tcctacgaag tgttcaagga tctcttcgac cccatcatcg
 361 aggaccggca cggcggctac aagcccagcg atgagcacaa gaccgacctc aaccccgaca
 421 acctgcaggg cggcgacgac ctggacccca actacgtgct gagctcgcgg gtgcgcacgg
 481 gccgcagcat ccgtggcttc tgcctccccc cgcactgcag ccgcggggag cgccgcgcca
 541 tcgagaagct cgcggtggaa gccctgtcca gcctggacgg cgacctggcg ggccgatact
 601 acgcgctcaa gagcatgacg gaggcggagc agcagcagct catcgacgac cacttcctct
 661 tcgacaagcc cgtgtcgccc ctgctgctgg cctcgggcat ggcccgcgac tggcccgacg
```

```
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 841 gcaccggcct cacccagatt gaaactctct tcaagtctaa ggactatgag ttcatgtgga
 901 accctcacct gggctacatc ctcacctgcc catccaacct gggcaccggg ctgcgggcag
 961 gtgtgcatat caagctgccc aacctgggca agcatgagaa gttctcggag gtgcttaagc
1021 ggctgcgact tcagaagcga ggcacaggcg gtgtggacac ggctgcggtg ggcggggtct
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1141 tggacggagt gaagctgctc atcgagatgg agcagcggct ggagcagggc caggccatcg
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1261 tcctaactta ttgcctgggc agtgcccacc atgcaccct gatgttcgcc gtctggcgag
1321 cccttagcct tgctgtagag acttccgtca cccttggtag agtttatttt tttgatggct
1381 aagatactgc tgatgctgaa ataaactagg gttttggcct gcctgcgtct g
```

SEQ ID NO. 35

X15334
Human gene for creatine kinase B (EC 2.7.3.2).

```
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  61 ttgcctgaag tctccgggga ctgcctttgg gggtcgggta aacttgtccc ctgcgaagag
 121 ggcccagggt tggggtctgg aaactccgag gctgcacttg ccagcggcct cttaaggcca
 181 cagcgtcccc gtggtttctg gctcgcagcc ccccgagacc caggacttgt ccaaggtcag
 241 ggcaccgcgg gtgcccccgg gctgggccgc agcagactgc gcttcccgcg cgccttcgct
 301 ttgcaccagg atcgcccagg aaatgcctgc gggcaccttg aggaaggtcg gcggctccgg
 361 gccagctcgc actggccggg gtggggcggg ggccgtacct gctgcggaag ccccgaaagc
 421 tttcgcccgg cccctcgccg ccgccgcggg ggctggctgg actaggcggg caggctcgag
 481 gatgcggatg aacccaagcg tcctcgagtg cccggaggct ctccgcctca gtttcccgcc
 541 cagaggcaag ggcgtgcgag gggatccaga tatccaagga cctgaggttt cggcctcgag
 601 gtcttgggcg ggggactggg caggctgcgc ggggtcccag cgaggggaca gctcgggtgg
 661 gcggccaggg tgttgggggc tgcgggcggc ggacaaagcg gcggcaccac cccgcggcgc
 721 gggccaatgg aatgaatggg ctataaatag ccgccaatgg gcggcccgcg ttgtgccccc
 781 taagagccgc gggagcgcgg agcggccgct gttcgcctgc gtcgctccgg gagctgccga
 841 cggacggagc gcccccgccc ccgcccggcc gcccggtgag tgggcccggg ggccgggggc
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 961 ggaccggccg ggggtccccg gcgatgatgg cgctccccgc gcgcgctgcg gaccccgctg
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1081 gggagcccgg gaaccccggc gtgccggtcc cctctgaccc cgcgtctccc cgcagcccgc
1141 cgccgccatg cccttctcca acagccacaa cgcactgaag ctgcgcttcc cggccgagga
1201 cgagttcccc gacctgagcg cccacaacaa ccacatggcc aaggtgctga cccccgagct
1261 gtacgcggag ctgcgcgcca agagcacgcc gagcggcttc acgctggacg acgtcatcca
1321 gacaggcgtg gacaacccgg gtacgcgacc cctcggggcc ggggtcccgg ccccccctcc
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1441 ccgtcccgcg ccccctcccc caggccaccc gtacatcatg accgtgggct gcgtggcggg
1501 cgacgaggag tcctacgaag tgttcaagga tctcttcgac cccatcatcg aggaccggca
1561 cggcggctac aagcccagcg atgagcacaa gaccgacctc aaccccgaca acctgcaggt
1621 gcggggctgc gggcgggccg ggcgggcggg gccggggtct tcgggcgctc actcccgtct
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1741 gggccgcagc atccgtggct tctgcctccc cccgcactgc agccgcgggg agcgccgagc
1801 catcgagaag ctcgcggtgg aaggtagggg ccgggcgggc cgaggggcgg cggcggccgc
1861 gtcccctcc cggcgcggtc cccgcccgct tttgtttacg tcgcccggga gcggcagccg
1921 ccgtcgcgct cttatctgcg cgcgcccggg ttcagtttcc cggacccacc gagggacgga
1981 ggcccagccc ccgcgcccac agcggcctgg ggcccaggga gggcgggtcc tggcgcgggg
2041 tcaccgcctg ggaccgtcgc ccgggccgtg aggactggac gcccgcagat ccgggcgggt
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```

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2161 ggaggggcgg gcgcgggagc ccagcgtcct gagcgcaccc ctcgcagccc tgtccagcct
2221 ggacggcgac ctggcgggcc gatactacgc gctcaagagc atgacggagg cggagcagca
2281 gcagctcatc gacgaccact tcctcttcga caagcccgtg tcgcccctgc tgctggcctc
2341 gggcatggcc cgcgactggc ccgacgcccg cggtatctgg tgcgtgtccc tctgcgccct
2401 ctcgcggcgt cctccctccc cgctacctcc gctttccctc tcgccccccc cgcgggggtg
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2581 gactgggccc cggcccgtgg gggctcagga ctcagcaaca cctccccacc ccgagacgtg
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2701 ttttcttttc tcactcctcc gccgctggga ttctaccagg ggctggtgac gccaaagctt
2761 ctccaggggc agggctccta cccccactgt gggggggcggg tcgggctgtc ctggcggtcc
2821 ctggccccgc cccacctcgg gccacagcgc atgatggcag ctggggttct cctgctgtga
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2941 ctcctgggca tggcagggag cagcacctca gggacttggg aagttccttt ggtctggggg
3001 cggcctgggg ctttttttctg ggtatgccct gagaccagcc ctcccgcagg cacaatgaca
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3961 gggttttggc ctgcctgcgt ctgagtggtg cctctccttt cccaggggg aggggaagg
4021 gcagcagcca ggccccagga gtcttgagtc ctgggcctgc tgtgggcctc gccttctgtg
4081 agatgggaca agagccagga ggtggccact ctgttctgcc tgccctacct agtccatggg
4141 ccccttccct cgtgtctatc gggctgtgca ggcaggaaca tgggagagag cgagggagga
```

SEQ ID NO. 36

P14618 - Homo sapiens
Pyruvate kinase M1 or M2 isozyme

```
mskphseagt afiqtqqlha amadtflehm crldidsppi tarntgiict igpasrsvet
lkemiksgmn varlnfshgt heyhaetikn vrtatesfas dpilyrpvav aldtkgpeir
tglikgsgta evelkkgatl kitldnayme kcdenilwld yknickvvev gskiyvddgl
islqvkqkga dflvteveng gslgskkgvn lpgaavdlpa vsekdiqdlk fgveqdvdmv
fasfirkasd vhevrkvlge kgknikiisk ienhegvrrf deileasdgi mvargdlgie
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awaedvdlrv nfamnvgkar gffkkgdvvi vltgwrpgsg ftntmrvvpv p
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SEQ ID NO. 37

X56494

H.sapiens M gene for M1-type and M2-type pyruvate kinase

```
   1 ggtcttcaca ttttgaatgc gcaacattgt atctgtgaat gaaggcaaga gttaacagct
  61 gtttaattga taactgctcg catcattagt tgctggctaa caactgggaa atcagaaaat
 121 gtcttgtaga aaaatgtaag aaaagttcca acaatactga cttaaacacg agcaaaggtg
 181 aaaacagaaa tgctgactcc tgcataggtt atcggccCta atgttctgac ttgatatttc
 241 cagatgccca gctctgcgct aatatcaaca ccgtctattt actttctact ctgaggcatt
 301 cgctctgcag gattccagac cctactaaat tattcacatg gccccaaccg gtccttcctt
 361 gttccgcggt cctaacacaa tgaatggtcc taagaggaaa acggcctcgg ctcccgctcc
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 481 aggcggccag gacctccaac cacgcacaag tctgcagctc tccccaactt tccgttcagc
 541 tcagtctccg agggtgcgcc agagcagaca cccggaggag tggggagtgg cagggcgggg
 601 ccgggagaat gctgccccgg aacccataaa ttcggccctg cccaggtagg ccgggacagc
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```

3181 agaagccccc gcccactcgg gctgaaggca gtgatgtggc caatgcagtc ctggatggag
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SEQ ID NO. 38

Q01995 - Homo sapiens
Transgelin

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SEQ ID NO. 39

D84342
Homo sapiens DNA for SM22 alpha, complete cds

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SEQ ID NO. 40

Q14103 - human
Heterogeneous nuclear ribonucleoprotein

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SEQ ID NO. 41

AF026126
Homo sapiens heterogeneous nuclear ribonucleoprotein D (HNRPD)
gene, complete cds

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SEQ ID NO. 42

NP_852000
GSK-3 Binding Protein - FRAT1

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SEQ ID NO. 43

NM_005479
Homo sapiens frequently rearranged in advanced T-cell lymphomas
FRAT1), transcript variant 1, mRNA

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SEQ ID NO. 44

NM_181355
Homo sapiens frequently rearranged in advanced T-cell lymphomas
FRAT1), transcript variant 2, mRNA

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1981 ttcatcatcc agatgtggct actgacatat ctacacttcg caccggagtg tctggaattg
2041 tggctatcct gattatagga ttttaactta actgaaatgc ctgctttgaa taaatgtgtt
2101 gggtttttg tttggtttta ttttatactt gccatcagtg aaaaagatgt acagaacaca
2161 tttctctgat ctccataaac atgaaaacac ttgaaatctc aaa
```

SEQ ID NO. 45

NP_444254
myosin light chain isform kinase 2

mgdvklvass hisktslsvd psrvdsmplt eapafilppr nlcikegata kfegrvrgyp
epqvtwhrng qpitsggrfl ldcgirgtfs lvihavheed rgkytceatn gsgarqvtve
ltvegsfakq lgqpvvsktl gdrfsasave trpsiwgecp pkfatklgrv vvkegqmgrf
sckitgrpqp qvtwlkgnvp lqpsarvsvs ekngmqvlei hgvnqddvgv ytclvvngsg
kasmsaelsi qgldsanrsf vretkatnsd vrkevtnvis keskldslea aakskncssp
qrggsppwaa nsqpqppres klesckdspr tapqtpvlqk tsssitlqaa rvqpeprapg
lgvlspsgee rkrpapprpa tfptrqpglg sqdvvskaan rripmegqrd safpkfeskp
qsqevkenqt vkfrceglav mevapsfssv lkdcaviegq dfvlqcsvrg tpvpritwll
ngqpiqyars tceagvaelh iqdalpedhg tytclaenal gqvscsawvt vhekkssrks
eyllpvapsk ptapiflqgl sdlkvmdgsq vtmtvqvsgn pppeviwlhn gneiqesedf
hfeqrgtqhs lciqevfped tgtytceawn sagevrtqav ltvqephdgt qpwfiskprs
vtaslgqsvl iscaiagdpf ptvhwlrdgk alckdtghfe vlqnedvftl vlkkvqpwha
gqyeillknr vgecscqvsl mlqnssaral prgrepasce dlcgggvgad gggsdrygsl
rpgwpargqg wleeedgedv rgvlkrrvet rqhteeairq qeveqldfrd llgkkvstkt
lseddlkeip aeqmdfranl qrqvkpktvs eeerkvhspq qvdfrsvlak kgtsktpvpe
kvpppkpatp dfrsvlggkk klpaengsss aetlnakave sskplsnaqp sgplkpvgna
kpaetlkpmg nakpaetlkp mgnakpdenl ksaskeelkk dvkndvnckr ghagttdnek
rsesqgtapa fkqklqdvhv aegkkllqc qvssdppati iwtlngktlk ttkfiilsqe
gslcsvsiek alpedrglyk cvakndagqa ecscqvtvdd apasentkap emksrrpkss
lppvlgtesd atvkkkpapk tppkaamppq iiqfpedqkv ragesvelfg kvtgtqpitc
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kpdppagtpc asdirssslt lswygssydg gsavqsysie iwdsanktwk elatcrstsf
nvqdllpdhe ykfrvrainv ygtsepsqes elttvgekpe epkdevevsd ddekepevdy
rtvtinteqk vsdfydieer lgsgkfgqvf rlvekktrkv wagkffkays akekenirqe
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segveyihkq givhldlkpe nimcvnktgt riklidfgla rrlenagslk vlfgtpefva
pevinyepig yatdmwsigv icyilvsgls pfmgdndnet lanvtsatwd fddeafdeis
ddakdfisnl lkkdmknrld ctqclqhpwl mkdtknmeak klskdrmkky marrkwqktg
navraigrls smamisglsg rksstgspts plnaeklese edvsqaflea vaeekphvkp
yfsktirdle vvegsaarfd ckiegypdpe vvwfkddqsi resrhfqidy dedgncslii
sdvcgdddak ytckavnslg eatctaeliv etmeegegeg eeeee

SEQ ID NO. 46

AF069601
Homo sapiens myosin light chain kinase isoform 2 (MLCK) mRNA, complete cds

ccggctgcct ctgctgcagt tcagagcaac ttcaggagct tcccagccga gagcttcagg
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ctatttctct caaaatcca

SEQ ID NO. 47

AAH07433 and P09493
tropomyosin 1 alpha chain.

1 mdaikkkmqm lkldkenald raeqaeadkk aaedrskqle delvslqkkl kgtedeldky
61 sealkdaqek lelaekkatd aeadvaslnr riqlveeeld raqerlatal qkleeaekaa
121 desergmkvi esraqkdeek meiqeiqlke akhiaedadr kyeevarklv iiesdlerae
181 eraelsegqv rqleeqlrim dqtlkalmaa edkysqkedr yeeeikvlsd klkeaetrae
241 faersvtkle ksiddledel yaqklkykai seeldhalnd mtsm

SEQ ID NO. 48

NM_000366 and BC007433
Homo sapiens tropomyosin 1 (alpha), mRNA (cDNA clone), complete cds

1 gaggaatgcg gtcgcccccc tgggaaagta catatctggg agaagcaggc ggctccgcgc
61 tcgcactccc gctcctccgc ccgaccgcgc gctcgccccg ccgctcctgc tgcagcccca
121 gggccccctcg ccgccgccac catggacgcc atcaagaaga agatgcagat gctgaagctc
181 gacaaggaga acgccttgga tcgagctgag caggcggagg ccgacaagaa ggcggcggaa
241 gacaggagca agcagctgga agatgagctg gtgtcactgc aaaagaaact caagggcacc
301 gaagatgaac tggacaaata ttctgaggct ctcaaagatg cccaggagaa gctggagctg

361 gcagagaaaa aggccaccga tgctgaagcc gacgtagctt ctctgaacag acgcatccag
421 ctggttgagg aagagttgga tcgtgcccag gagcgtctgg caacagcttt gcagaagctg
481 gaggaagctg agaaggcagc agatgagagt gagagaggca tgaaagtcat tgagagtcga
541 gcccaaaaag atgaagaaaa aatggaaatt caggagatcc aactgaaaga ggcaaagcac
601 attgctgaag atgccgaccg caaatatgaa gaggtggccc gtaagctggt catcattgag
661 agcgacctgg aacgtgcaga ggagcgggct gagctctcag aaggccaagt ccgacagctg
721 gaagaacaat taagaataat ggatcagacc ttgaaagcat taatggctgc agaggataag
781 tactcgcaga aggaagacag atatgaggaa gagatcaagg tcctttccga caagctgaag
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901 gatgacttag aagacgagct gtacgctcag aaactgaagt acaaagccat cagcgaggag
961 ctggaccacg ctctcaacga tatgacttcc atgtaaacgt tcatccactc tgcctgctta
1021 caccctgccc tcatgctaat ataagtttct ttgcttcact tctcccaaga ctccctcgtc
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1141 ctctctctta gcatcctgcc ttagagccag gcacacactg tgctttctat tgtacagaag
1201 ctcttcgttt cagtgtcaaa taaacactgt gtaagctaaa aaaaaaaaaa aaaaaa

SEQ ID NO. 49

EITALAPSTMK

SEQ ID NO. 50

MLTELEK

SEQ ID NO. 51

ALNSIIDVYHK

SEQ ID NO. 52

GADVWFK